SEQUENCE LISTING

- (1) GENERAL INFORMATION:
 - (i) APPLICANT: Thomson, James
 - (ii) TITLE OF INVENTION:

Primate Embryonic Stem Cells

- (ii\i) NUMBER OF SEQUENCES: 6
 - (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Quarles & Brady
 - (B) STREET: 411 East Wisconsin Avenue
 - (C) CITY: Milwaukee
 - (D) STATE: WI
 - (E) COUNTRY: US
 - (F) ZIP:\53202
 - (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: \\TBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: Pate tin Release #1.0,

Version #1.25

- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER:
 - (B) FILING DATE:
 - (C) CLASSIFICATION:
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Schwartz, Carl R.
 - (B) REGISTRATION NUMBER: 29,437
 - (C) REFERENCE/DOCKET NUMBER: 960296\9540-8
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: (414) 277-5715
 - (B) TELEFAX: (414) 271-3552

(2) INFORMATION FOR SEQ ID NO:1:	
(i) SEQUENCE CHARACTERISTICS:	
(A) LENGTH: 25 base pairs	
(B) TYPE: nucleic acid	
(C) STRANDEDNESS: double	
(D) TOPOLOGY: linear	
(\daggeright) MOLECULE TYPE: DNA (genomic)	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:	
GCTGGATTOT CTGCAGGATG GGGAA	25
(2) INFORMATION FOR SEQ ID NO:2:	
(i) SEQUENCE CHARACTERISTICS:	
(A)\LENGTH: 25 base pairs	
(B) TYPE: nucleic acid	
(C) STRANDEDNESS: double	
(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: DNA (genomic)	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:	
TCCCCTGAAG AAAATTGGTT AAAAT	25
(2) INFORMATION FOR SEQ ID NO:3:	
(i) SEQUENCE CHARACTERISTICS:	
(A) LENGTH: 29 base pairs	
(B) TYPE: nucleic acid	
(C) STRANDEDNESS: double	
(D) TOPOLOGY: linear \	
(ii) MOLECULE TYPE: DNA (genomic)	
(xi) SEQUENCE DESCRIPTION: SEQ 10 NO:3:	
GGATCCACCG TCAACACCAC CATCTGTGC	29
(2) INFORMATION FOR SEQ ID NO:4:	
(i) SEQUENCE CHARACTERISTICS:	
(A) LENGTH: 30 base pairs	
(B) TYPE: nucleic acid	

(C) STRANDEDNESS: double	
(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: DNA (genomic)	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:	
GGATCCACAG GTCAAAGGGT GGTCCTTGGG 3	0
(2) INFORMATION FOR SEQ ID NO:5:	
(i) SEQUENCE CHARACTERISTICS:	
(A) LENGTH: 30 base pairs	
(B) TYPE: nucleic acid	
(C) STRANDEDNESS: double	
(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: DNA (genomic)	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:	
GGGAATTCGC AGTTACTGAG AAGTCACAAG 3	0
(2) INFORMATION FOR SEQ ID NO:6:	
(i) SEQUENCE CHARACTER STICS:	
(A) LENGTH: 30 base pairs	
(B) TYPE: nucleic adid	
(C) STRANDEDNESS: double	
(D) TOPOLOGY: linear \	
(ii) MOLECULE TYPE: DNA (genomic)	
(xi) SEQUENCE DESCRIPTION: SEQ D NO:6:	
GGGAATTCGA AGCATGTCAA AGTGGTATGG \	0
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